Resource Summary Report

Generated by PRECISE-TBI on Sep 7, 2026

CopyDetective

RRID:SCR_018909
Type: Tool

Proper Citation

CopyDetective (RRID:SCR_018909)

Resource Information

URL: https://github.com/sandmanns/CopyDetective

Proper Citation: CopyDetective (RRID:SCR_018909)

Description: Software tool for detection threshold aware CNV calling in matched whole exome sequencing data.

Resource Type: algorithm resource, data analysis software, data processing software, software application, software resource

Keywords: Copy number variant calling, whole exome sequencing data, detection thresholds, deletion, duplication, case control samples, bio.tools

Funding: DFG TU 298/5-1 (DFG Clinical Research Unit 326 Male Germ Cells: from Genes to Function);
DKH 111347;
DKS DKS349 2014.11 A/B (NHL-BFM Registry 2012);
EU 634789 (Horizon2020 MDS-RIGHT);
Löwenkinder - Verein zur Unterstützung krebskranker Kinder e.V.

Availability: Free, Available for download, Freely available

Resource Name: CopyDetective

Resource ID: SCR_018909

Alternate IDs: biotools:copydetective

Alternate URLs: https://bio.tools/copydetective

License: GNU Affero General Public License v3.0

Record Creation Time: 20220129T080342+0000

Record Last Update: 20260905T132843+0000

Ratings and Alerts

No rating or validation information has been found for CopyDetective.

No alerts have been found for CopyDetective.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at PRECISE-TBI .

Sandmann S, et al. (2020) CopyDetective: Detection threshold-aware copy number variant calling in whole-exome sequencing data. GigaScience, 9(11).